

DATE: 02726/2003

1600

## PATENT APPLICATION: US/09/804,733C TIME: 14:57:02 Input Set : D:\Mons016.app Output Set: N:\CRF4\02262003\I804733C.raw 3 <110 · APPLICANT: WANG, QI ET AL. 5 - 120 - TITLE OF INVENTION: RECOMBINANT PROTEINS CONTAINING REPEATING UNITS T 4:130 FILE FEFERENCE: MONS:016US 9 k140 CUFRENT APPLICATION NUMBER: 09/804,733C C--> 10 <141> CURRENT FILING DATE: 2001-03-13 Does No: Comply 12 H156 - PRIOR APPLICATION NUMBER: PCT/US01/07957 Corrected Diskette Needec 13 -(151 > PFIOR FILING DATE: 2001-03-13 15 - 160 - NUMBER OF SEQ ID NOS: 29 17 <170 - SOFTWARE: PatentIn Ver. 2.1 19 K210% SEQ ID NO: 1 20 4211 > LENGTH: 5 01 KO1D: TYPE: PRT 11 +213 + OFGANISM: Euthynnus pelamis 14 4400 / SEQUENCE: 1 15 Leu Lys Pro Ash Met 19 - 3105 SEQ ID NO: 2 30 - 211 - LENGTH: 4 31 4212 - TYFE: FRT 32 (313 - OFGANISM: Euthynnus pelamis 34 -400 - SEQUENCE: 2 35 Lys Pro Ash Met 3.6 39 H210H SEQ IF NO: 3 40 HILLS LENGTH: 4 41 KEIDS TYPE: PRT 40 0213 POFGANISM: Euthynnus pelamis 44 < 400 > SEQUENCE: 3 45 Val Val Tyr Ero 1 46 49 -210 - SEQ ID NO: 4 50 k211 + MENGTH: 15 51 - Clus TYPE: DNA 52 4213 CFGANISM: Artificial Sequence 54 HODD - FEATURE: 55 - 2003 - OTHER INFOFMATION: Description of Artificial Sequence: Synthetic C 15 Primer 58 -0020 - FEATURE: 59 -0000 NAME/KEY: modified\_base #60 <2222 + LCCATION: (3)..(3) 61 $\pm 223$ OTHER INFORMATION: N = A, C, G or T/U 63 HAUL - SEQUENCE: 4 15

RAW SEQUENCE LISTING

W--> 64 ctnaarccna ayatg

init: ....

TIME: 14:67:57

Input Set : D:\Mons016.app Cutcut Set: N:\CRF4\02262003\I804733C.raw 67 k210% SEQ ID NO: 5 68 <311 - LENGTH: +-0 69 K212 · TYFE: DNA 73 (1813 - ORGANISM: Artificial Sequence 71 : 122 1: FLATURE: 7: 322 OTHER INFORMATION: Description of Artificial Sequence: Synthetic 7.4 Frimer 7: K22 : FEATUPE: 77 HARL- NAME/FEY: modified base 78 -: 2225 LOCATION: (3)..(54) 79 + 20 N + OTHER INFORMATION: N = A, C, G, cr T/U81 KADOW SEQUENCE: 5 W--> 82 ctnaarccna ayatgctnaa rccnaayatg ctnaarccna ayatgctnaa rccnaayatg 60 85 -2200 SEQ II NO: € 86 -1111: LENGTH: 60 87 GLILL TYPE: DNA 86 - MILTO OFGANISM: Artificial Sequence 90 FILLOR FEATURE: 91 KARAS OTHER INFORMATION: Description of Artificial Sequence: Synthetic 91 Primer 94 -COCS FEATURE: 95 W221> NAME/KEY: modified base 96 (1.11): LOCATION: (7). ((52)) (53) 97 +0.23 OTHER INFORMATION: N = A, C, G or T/U 99 4400 SEQUENCE: 6 W--> 100 catrttnggy ttnagcatrt tnggyttnag catrttnggy ttnagcatrt tnggyttnag 60 103 K0105 SEQ ID NO: 7 164 K211 - LENGTH: 25 105 KIII TYPE: DNA 10% HULLS ORGANISM: Artificial Sequence 10% HOSO - FEATURE: 109 - 215 - OTHER INFORMATION: Description of Artificial Sequence: Synthetic Primer 110 110 k010 - FEATURE: 113 KODA'S NAME/KEY: modified base 114 <2220 LOCATION: (12)..(18) 115 <0.130 OTHER INFORMATION: N = A, C, G or T/U 117 <400 > SEQUENCE: 7 25 W--> 118 aaagaattcc tnaarccnaa yatgc 1.1 <3105 SEQ ID NO: 8 103 K010x TYPE: DNA 124 <213> ORGANISM: Artificial Sequence 126 KOBO - FEATURE: 127 - COMES OTHER INFORMATION: Description of Artificial Sequence: Synthetic 125 Primer 130 KIDC - FEATURE: 131 - MAME/KEY: modified base

RAW SEQUENCE LISTING

FATENT AFFLICATION: US/09/804,733C

131 H2227 LOCATION: (18)..(24)



## RAW SEQUENCE LISTING

FATERT AFFILICATION: US/09/804,733C

TATE: 42 is 7000 TIME: 14:87:32

27

20

Input Set : D:\Mons016.app

Output Set: N:\CRF4\02262003\I804733C.raw

133 <223 > OTHER INFORMATION: N = A, C, G or T/U

135 -400 - SEQUENCE: 8

# W--> 136 aaagcggccg ccatrttngg yttnagc

SED ID NO: 3 139 +.11 SED ID NO: 143 +/11 + DENGTH: 20

141 - 11 - TYPE: DNA

140 - 133 - CR3ANISM: Artificial Sequênce

144 - JOH - FEATUFE:

145 - LL: OTHER INFOFMATION: Description of Artificial Sequence: Synthetic

146 Frimer

148 4440 + SEQUENCE: 9

149 taatacgast cactataggg

15.1 - 210.4 SEQ IE NO: 10

153 - 211 - LENGTH: 19

154 - 212 - TYPE: DNA

155 - 113 - ORGANISM: Artificial Sequence

157 SILON FEATURE:

198 - 118 - OTHER INFORMATION: Description of Artificial Sequence: Synthetic

Primer 159

161 < 400 > SEQUENCE: 10

19 160 ogatomatam ogagrogod

165 - 210% SEQ ID NO: 11

166 - 211 - LENGTH: 45

167 k212% TYPE: DNA

168 klife ORGANISM: Artificial Sequence

170 - 100 - FEATUFE:

171 42.30 OTHER INFOFMATION: Description of Artificial Sequence: Synthetic

Frimer

174 < 120 + FEATUFE:

175 - III - MAME/KEY: rediffied\_base

176 - 222 - LOCATION: (3)..(48)

177 - LIE - OTHER INFOFMATION: N = A, C, G or T/U

179 - 400 - SEQUENCE: 11

#### 48 W--> 180 gtngtntayc cngtngtnta yccngtngtn tayccngtng tntayccn

193 k2105 SEQ ID NO: 12

194 - 211 - LENGTH: 48

185 KIII TYPE: DNA

186 - J13 - OEGANISM: Artificial Sequence

188 /2200 FEATURE:

189 < 2235 OTHER INFORMATION: Description of Artificial Sequence: Synthetic

190 Prime:

191 <210 · FEATURE:

193 <221 NAME/KEY: modified\_base

194 \*222\* LOCATION: (1)..(46)

195 <223 OTHER INFORMATION: N = A, C, G or T/U

197 - 400 - SEQUENCE: 12

#### 48 W--> 198 nggrtanach acnggrtana chachggrta nachachggr tanachac

201 - 110 - SEQ ID NO: 13

202 · J11 · LENGTH: 33

TATE: ... president

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TIME: 14:6": 2
                     PATENT APPLICATION: US/09/804,733C
                     Input Set : D:\Mons016.app
                     Output Set: N:\CRF4\02262003\1804733C.raw
     203 - Lif - TYPE: DNA
     264 - 213 - ORGANISM: Artificial Sequence
     206 (.L. FEATURE:
     197 ... CTHER INFORMATION: Description of Artificial Sequence: Synthetic
              Frimer
     ::0 · LL. · FEATUFE:
     ::: ...: NAME/FEY: modified_base
     111 - 111 - 16 TATION: (12)...33)
     113 - 113 - CIHEP INFORMATION: N = A, C, G or T/U
     33
W--> 216 aaaggatccg tngtntaycc ngtngtntay ccn
     119 - 110 - SEQ II NO: 14
     120 -011 - LENGTH: 33
     Duly (1912) TYPE: INA
     722 - 213 - OF GANISM: Artificial Sequence
     .114 ×12.00 FEATUFE:
     215 WILLS OTHER INFOFMATION: Description of Artificial Sequence: Synthetic
              Frimer
     DIS KODO - FEATUFE:
     P29 - 211 - NAME/KEY: mcdified_base
     230 KISI - LOCATION: (10)..(31)
     23: +.23 + 0THER INFORMATION: N = A, C, G or T/U
     223 k400 > SEQUENCE: 14
                                                                               33
W--> 234 cccaagettn ggrtanacna enggrtanac nac
     337 ×210> SEQ ID NO: 15
     238 V.111 - LENGTH: 45
     130 - M11 - TYPE: DNA
     140 - 1130 OFGANISM: Artifleial Sequence
     140 KILON FEATURE:
     143 - 123 - OTHER INFORMATION: Description of Artificial Sequence: Synthetic
     Frimer
     746 × 2200 FEATURE:
     [47 . Dits MAME (REY: modified_base
     248 ×200 × LOCATION: (3)..(45)
     049 \times 103 \times \text{OTHER INFOPMATION: N = A, C, G or T/U}
     251 K4005 SEQUENCE: 15
W--> 252 gtnceneeng theeneengt neeneength ceneengthe encen
                                                                               45
     155 k210% SEQ ID NO: 16
      356 <2115 LENGTH: 45
      357 HOIDH TYEE: DNA
     158 - 213 - OFGANISM: Artificial Sequence
     160 KIIOV FEATURE:
     261 4223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
     262
               Primer
     364 - 200 - FEATUPE:
     165 + 121 > NAME/KEY: modified_base
      266 SELES LOCATION: (1)..(43)
     167 \cdot 113 \cdot \text{OTHER INFORMATION: N = A, C, G or T/U}
     269 -450% SEQUENCE: 16
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RAW SEQUENCE LISTING

TIME: 14:87:10

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Input Set : D:\Mons016.app
                     Output Set: N:\CRF4\02262003\1804733C.raw
                                                                              45
W--> 270 nggnggnacn ggnggnacng gnggnacngg nggnacnggn ggnac
     273 +010 - SEQ II NO: 13
     274 - 11 - LENGTE: 36
     275 - 31. - TYFE: DNA
            278 · J. J · FEATURE:
     279 - 1. : - CIHER INFORMATION: Description of Artificial Sequence: Synthetic
              Frim∈r
     297 - ... FEATURE:
     LEF + DLT + NAME, FEY: modified_base
     L84 \times .12.1 \times 1.00 \text{ PATION: } (12) \dots (36)
     LES - DIB - GIREF INFOFMATION: N = A, C, G or T/U
     28% - 40% - SEQUENCE: 17
                                                                              36
W--> 288 aaaggateeg tneeneengt neeneengtn eeneen
     191 - 110 - SEQ ID NO: 18
     192 - 211 - LENGTH: 36
     193 KININ TYPE: CNA
     194 4213 + OFGANISM: Artificial Sequence
      196 -210 - FEATUFE:
     197 (173) OTHER INFORMATION: Description of Artificial Sequence: Synthetic
     <u> 198</u>
               Frimer
     300 + 320 + FEATUFE:
     PG1 + 101 + NAME/KEY: mcdified_base
     FOR RESENT LOCATION: (10)..(34)
     503 + 203 + OTHEF INFOFMATION: N = A, C, G or T/U
     305 - 400 - SEQUENCE: 18
                                                                               36
W--> 306 aataagcttn ggnggnacng gnggnacngg nggnac
      909 - 210 - SEQ ID NO: 19
      311 - LIL - TYFE: PET
      310 - 213 - OFGANISM: Artificial Sequence
      314 + 200 + FEATURE:
      315 0203 OTHER INFOFMATION: Description of Artificial Sequence:m Synthetic
      35.6
               Feptide
      318 K400% SEQUENCE: 13
      319 Val Pro Pro Leu Lys Pro Asn Met
      320
      303 k010 SEQ ID NO: 10
      304 <0115 LENGTH: 48
      BD5 - DID - TYPE: DNA
      3.6 *213 * OFGANISM: Artificial Sequence
      328 - 220 - FEATURE:
      329 8223 OTHER INFORMATION: Description of Artificial Sequence: Synthetic
      330
               Frimer
      930 WELLON FRATURE:
      333 k231 + NAME/KEY: modified_base
      334 kD12+ LOCATION: (3)..(42)
      135 < 0.35 OTHER INFORMATION: N = A, C, G or T/U
      337 - 400 > SEQUENCE: 20
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RAW SEQUENCE LISTING

FATENT APPLIBATION: US/09/804,733C

RAW SEQUENCE LISTING ERROR SUMMARY FATERIT APPLICATION: US/09/804,733C

TIME: 14:50:03

Input Set : D:\Mons016.app

Output Set: N:\CRF4\02262003\I804733C.raw

### Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seg#:4; N Pos. 3,9 Seq#:5; N Fos. 3,9,18,24,33,39,48,54 Seq#:6; N Pos. 7,13,22,28,37,43,52,58 Seq#:"; N Pos. 12,18 Seq#:8; N Pcs. 18,24 Seq#:11; N Fcs. 3,6,12,15,18,24,27,30,36,39,42,48 Seq#:12; N Fos. 1,7,10,13,19,22,25,31,34,37,43,46 Seq#:13; N Pos. 12,15,21,04,07,33 Seq#:14; N Pas. 10,16,19,22,28,31 Seq#:15; N Pos. 3,6,9,12,15,18,11,24,27,30,33,36,39,42,45 Seq#:10; N Pos. 1,4,7,10,13,16,19,2%,25,18,31,34,37,40,43 Seq#:17; N Pos. 12,715,18,31,34,17,30,33,36 Seq#:18; N Pcs. 10,13,16,13,22,25,24,31,34 Seq#:20; N Pcs. 3,6,9,12,18,27,30,33,36,42 Seq#:21; N Pcs. 7,13,16,14,22,31,37,40,43,46 Seq#:22; N Pcs. 13,16,14,22,31,37,40,43,46,52 Seq#:23; N Pas. 19,25,29,31,34,40,43,46,85,61,64,67,70,76,79,82 Seq#:05; 11 Pcs. 3,6,9,15,18,.1,14,30,39,41,45,51,54,57,60,66 Seq#:26; N Pos. 7,13,16,19,20,20,31,34,43,49,52,55,58,64,67,70 Seq#:27; N Pos. 13,16,19,85,28,31,34,40,49,52,55,61,64,67,70,76 Seq#:28; N Pos. 19,25,28,31,34,40,43,46,55,61,64,67,70,76,79,82